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Timestamp: [year=2009; month=7; day=24; hr=11; min=17; sec=9; ms=277; ]

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Application No: 10573372 Version No: 4.0

Input Set:

Output Set:

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Finished: 2009-07-14 14:14:38.535  
Elapsed: 0 hr(s) 0 min(s) 5 sec(s) 991 ms  
Total Warnings: 3  
Total Errors: 0  
No. of SeqIDs Defined: 7  
Actual SeqID Count: 7

Error code	Error Description
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<110> Dale, James Langham  
Echeverria, Santy Peraza

<130> DAVI172.006APC

<141> 2009-07-14

<151> 2004-09-23

<151> 2003-09-25

<170> FastSEQ for Windows Version 4.0

<213> Musa acuminata

 $\langle 222 \rangle \quad (1) \dots (4323)$ 

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Gln Thr Leu Leu Asp Lys Ala Ser Asn Cys Ala Ile Gln Gln Leu Ala  
20 25 30

cgc cgc cgc ggc ctt cac gat gac ctg agg cgc ctg cgg acg tct ctg 144  
Arg Arg Arg Gly Leu His Asp Asp Leu Arg Arg Leu Arg Thr Ser Leu  
35 40 45

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50 55 60

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65 70 75 80

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Tyr Asp Ala Glu Asp Leu Leu Glu Glu Leu Glu Tyr Gln Ala Ala Lys	
85 90 95	
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Gln Lys Val Glu His Arg Gly Asp Gln Ile Ser Asp Leu Phe Ser Phe	
100 105 110	
tcc ctt agt act gcg agc gag tgg ttg ggt gcc gat ggt gat gat gct	384
Ser Leu Ser Thr Ala Ser Glu Trp Leu Gly Ala Asp Gly Asp Asp Ala	
115 120 125	
ggg act cga ttg agg gag atc cag ggg aag ctg tgc aac att gct gcc	432
Gly Thr Arg Leu Arg Glu Ile Gln Gly Lys Leu Cys Asn Ile Ala Ala	
130 135 140	
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145 150 155 160	
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Gln Phe Asp Trp Lys Val Val Arg Arg Glu Thr Ser Ser Phe Leu Thr	
165 170 175	
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Glu Thr Val Val Phe Gly Arg Asp Gln Glu Arg Glu Lys Val Val Glu	
180 185 190	
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Tyr Asn Asp Asn Arg Val Gly Asn Tyr Phe His Leu Lys Val Trp Val	
225 230 235 240	
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Cys Val Ser Asp Asn Phe Asn Val Lys Arg Leu Thr Lys Glu Ile Ile	
245 250 255	
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Glu Ser Ala Thr Lys Val Glu Gln Ser Asp Lys Leu Asn Leu Asp Thr	
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Leu Gln Gln Ile Leu Lys Glu Lys Ile Ala Ser Glu Arg Phe Leu Leu	
275 280 285	
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Val Leu Asp Asp Val Trp Ser Glu Asn Arg Asp Asp Trp Glu Arg Leu	
290 295 300	
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Cys Ala Pro Leu Arg Phe Ala Ala Arg Gly Ser Lys Val Ile Val Thr	

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act cga gac aca aag att gcc agc atc att ggc aca atg aag gaa att				1008
Thr Arg Asp Thr Lys Ile Ala Ser Ile Ile Gly Thr Met Lys Glu Ile				
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Ser Leu Asp Gly Leu Gln Asp Asp Ala Tyr Trp Glu Leu Phe Lys Lys				
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Cys Ala Phe Gly Ser Val Asn Pro Gln Glu His Leu Glu Leu Glu Val				
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Ile Gly Arg Lys Ile Ala Gly Lys Leu Lys Gly Ser Pro Leu Ala Ala				
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Lys Thr Leu Gly Ser Leu Leu Arg Leu Asp Val Ser Gln Glu His Trp				
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Leu Arg Gln Cys Phe Ala Phe Cys Ala Val Phe His Lys Asp Tyr Leu				
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Phe Tyr Lys His Glu Leu Ile Gln Thr Trp Ile Ala Glu Gly Phe Ile				
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gca cat caa gga aac aag agg atg gaa gat gtc gga agc agc tac ttc				1440
Ala His Gln Gly Asn Lys Arg Met Glu Asp Val Gly Ser Ser Tyr Phe				
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cat gag ctt gtt aat agg tct ttc ttt cag gaa tct cgg tgg aga ggg				1488
His Glu Leu Val Asn Arg Ser Phe Phe Gln Glu Ser Arg Trp Arg Gly				
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cga tat gtg atg cat gac ctc ata cac gat ctt gcc caa ttt ata tca				1536
Arg Tyr Val Met His Asp Leu Ile His Asp Leu Ala Gln Phe Ile Ser				
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Val Gly Glu Cys His Arg Ile Asp Asp Asp Lys Ser Lys Glu Thr Pro				
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Ser Thr Thr Arg His Leu Ser Val Ala Leu Thr Glu Gln Met Lys Leu				
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Gln	Arg	Asn	Gln	Tyr	Pro	Tyr	Met	Thr	Lys	Val	Asn	Ser	Cys	Leu	Leu	
565			570			575										
cct	cat	agc	ttg	ttc	aaa	aga	ctg	aaa	aga	atc	cat	gtt	tta	gtt	ttg	1776
Pro	His	Ser	Leu	Phe	Lys	Arg	Leu	Lys	Arg	Ile	His	Val	Leu	Val	Leu	
580			585			590										
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Gln	Leu	Arg	Tyr	Leu	Asp	Ile	Ser	Tyr	Asn	Ala	Cys	Ile	Gln	Arg	Leu	
610			615			620										
ccc	gag	tca	ttg	tgc	gac	ctt	tac	aat	ctg	caa	gca	ctg	agg	cta	tgg	1920
Pro	Glu	Ser	Leu	Cys	Asp	Leu	Tyr	Asn	Leu	Gln	Ala	Leu	Arg	Leu	Trp	
625			630			635										
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Gly	Cys	Gln	Leu	Arg	Ser	Phe	Pro	Gln	Gly	Met	Ser	Lys	Leu	Ile	Asn	
645			650			655										
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675			680			685										
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Glu	Ala	Ser	Lys	Ala	Lys	Leu	His	Arg	Lys	Gln	Tyr	Leu	Glu	Ala	Leu	
725			730			735										
gag	tta	gag	tgg	gca	gct	ggc	cag	gtt	tcc	agc	ttg	gag	cat	gag	tta	2256
Glu	Leu	Glu	Trp	Ala	Ala	Gly	Gln	Val	Ser	Ser	Leu	Glu	His	Glu	Leu	
740			745			750										
ctt	gtc	tcg	gag	gaa	gta	ctt	tta	ggg	ctc	caa	cca	cat	cac	ttc	ctc	2304
Leu	Val	Ser	Glu	Glu	Val	Leu	Leu	Gly	Leu	Gln	Pro	His	His	Phe	Leu	
755			760			765										

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785 790 795 800	
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Cys Thr Arg Leu Glu Gly Leu Ser Tyr Ile Gly Gln Leu Pro His Leu	
805 810 815	
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Lys Val Leu His Met Lys Arg Met Pro Val Val Lys Gln Met Ser His	
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Leu Val Leu Glu Asp Met Pro Thr Leu Lys Glu Phe Pro Asn Leu Ala	
850 855 860	
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Gln Leu Pro Cys Leu Lys Ile Ile His Met Lys Asn Met Phe Ala Val	
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Leu Ser Leu Glu Glu Leu Val Leu Gln Asp Met Leu Thr Leu Glu Glu	
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Leu Pro Asn Leu Gly Gln Leu Pro His Leu Lys Val Ile His Met Lys	
915 920 925	
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Leu	Thr	Phe	Glu	Glu	Leu	His	Ser	Leu	Glu	Lys	Leu	Pro	Cys	Leu		
1010		1015					1020									
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Lys	Val	Phe	Arg	Ile	Lys	Gly	Leu	Pro	Ala	Val	Lys	Lys	Ile	Gly		
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Leu	Glu	Asp	Leu	Val	Leu	Ser	Asp	Met	Pro	Ala	Trp	Glu	Glu	Trp		
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Ser	Trp	Ala	Glu	Arg	Glu	Glu	Leu	Phe	Ser	Cys	Leu	Cys	Arg	Leu		
1070		1075					1080									
aaa	att	gaa	caa	tgc	ccc	aaa	ctt	aaa	tgc	ttg	ctt	ccc	atc	cct		3294
Lys	Ile	Glu	Gln	Cys	Pro	Lys	Leu	Lys	Cys	Leu	Leu	Pro	Ile	Pro		
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1100		1105					1110									
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Thr	Ala	Ser	Leu	Ser	Leu	Leu	His	Ile	Ile	Lys	Cys	Pro	Asn	Leu		
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Arg	Asn	Leu	Gly	Glu	Gly	Leu	Leu	Ser	Asn	His	Leu	Pro	His	Ile		
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Asn	Ala	Ile	Arg	Ile	Trp	Glu	Cys	Ala	Glu	Leu	Leu	Trp	Leu	Pro		
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Val	Lys	Arg	Phe	Arg	Glu	Phe	Thr	Thr	Leu	Glu	Asn	Leu	Ser	Ile		
1175		1180					1185									
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Arg	Asn	Cys	Pro	Lys	Leu	Met	Ser	Met	Thr	Gln	Cys	Glu	Glu	Asn		
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gac	ctc	ctc	ctc	ccg	ccg	tta	atc	aag	gca	cta	gaa	ttg	ggt	gac		3654
Asp	Leu	Leu	Leu	Pro	Pro	Leu	Ile	Lys	Ala	Leu	Glu	Leu	Gly	Asp		



1205

1210

1215

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